

SEQUENCE LISTING

<110> Caskey, C. Thomas
Shumaker, John
Metspalu, Andres

<120> PARALLEL PRIMER EXTENSION APPROACH TO
NUCLEIC ACID SEQUENCE ANALYSIS

<130> 2875.1001-007

<150> US 08/564,100

<151> 1994-06-22

<150> SE 9302152-5

<151> 1993-06-22

<160> 22

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> polynucleotide of interest

<400> 1

gatagcaatc gcttacggta atccggcctg

30

<210> 2

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

<400> 2

gatagcaatc

10

<210> 3

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

<400> 3

atagcaatcg

10

<210> 4

<211> 10

<212> DNA

<223> oligonucleotide primer

10

<220>
<223> Oligonucleotide primer

10

10

10

10

10

<211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Oligonucleotide primer

<400> 15
 tgccattagg 10

<210> 16
 <211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Oligonucleotide primer

<400> 16
 gccattaggc 10

<210> 17
 <211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Oligonucleotide primer

<400> 17
 ccattaggcc 10

<210> 18
 <211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Oligonucleotide primer

<400> 18
 cattaggccg 10

<210> 19
 <211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Oligonucleotide primer

<400> 19
 attaggccgg 10

<210> 20
 <211> 10
 <212> DNA
 <213> Artificial Sequence

Sequence

10

10

30